

PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/763,011

DATE: 03/02/2001
TIME: 12:57:28

Input Set : A:\CANDIDA.txt
Output Set: N:\CRF3\03022001\I763011.raw

Does Not Comply
Corrected Diskette Needed

pp 6-7

OK

5 <110> APPLICANT: Contreras, Roland
6 Nelissen, Bart
7 DeBacker, Marianne
8 Luyten, Walter
9 Viaene, Jasmine
10 Logghe, Marc George
13 <120> TITLE OF INVENTION: Drug Targets In Candida Albicans
16 <130> FILE REFERENCE: 50899/002
19 <140> CURRENT APPLICATION NUMBER: US/09/763,011
21 <141> CURRENT FILING DATE: 2001-02-14
24 <150> PRIOR APPLICATION NUMBER: 98310694.9
26 <151> PRIOR FILING DATE: 1998-12-23
29 <150> PRIOR APPLICATION NUMBER: 9817796.7
31 <151> PRIOR FILING DATE: 1998-08-14
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66 gtcggtgata tgtacggtag aagagttggt atcatgactg ctatgatcat ttatattggt 420
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134 aattcaattc caatatggtc atgacaattg aaagtgttac tgtctgtgtt tacatattct 180
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142 ggttgatatc attgactaaa actaccctag ggataaatgc tgaaccgtgg ttaccaactg 420
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202 tacgaagatg atggtaaaga agacgatagt gatgagatta cggctcaaat actattatcc 1140
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206 gaagacaata aagtaatatc tgtaataaca ccaaagacaa agaaacctac tacaacagta 1260
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238 20 25 30
242 Ile Ser Ile Leu Thr Asn His Phe Gln Ile Leu Lys Asp Leu Leu Pro
244 35 40 45
248 Tyr Ser Lys Thr Ser Lys Pro Gln Ile Lys Glu Ser Arg Pro Leu Ile
250 50 55 60
254 Lys Val Ser Arg Asp Gly Val Pro Ile Asn Phe His Arg Ala Pro Ala
256 65 70 75 80
260 Ile Ile Met Lys Ser Asn Lys Thr Asp Asp Leu Val Arg Asn Ser Asn
262 85 90 95
266 Lys Thr Met Val Leu Thr Glu Ile Lys Thr Ile Thr Glu Phe Ala Thr
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274 115 120 125
278 Asn Thr Leu Ser Ile Glu Thr Ser Thr Pro Thr Phe Gln Ser His Asp
280 130 135 140
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286 145 150 155 160
290 Glu Ser Ser Asp Ala Leu Gln Arg Asp Ala Phe Asp Gln Ile Lys Lys
292 165 170 175
296 Leu Glu Lys Leu Val Leu Asp Leu Arg Leu Glu Met Lys Glu Gln Gln
298 180 185 190
302 Lys Ser Phe Asn Asp Gln Leu Val Asp Ile Tyr Thr Ala Arg Ser Ile
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308 Val Pro Ile Tyr Thr Thr His Ile Val Thr Ser Ala Ile Pro Ser Tyr
310 210 215 220
314 Val Pro Lys Glu Glu Val Met Val Ser His Asp Thr Ala Pro Ile Val
316 225 230 235 240
320 Ser Arg Pro Arg Thr Asp Ile Pro Val Ser Gln Arg Ile Asp Thr Ile
322 245 250 255
326 Ser Lys His Lys Met Asn Gly Lys Asn Ile Leu Asn Asn Asn Pro Pro
328 260 265 270
332 Pro Asn Ser Val Leu Ile Val Pro Gln Phe Gln Phe His Glu Arg Met
334 275 280 285
338 Ala Thr Lys Thr Glu Val Ala Tyr Met Lys Pro Lys Ile Val Trp Thr
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344 Asn Phe Pro Thr Thr Thr Ala Thr Ser Met Phe Asp Asn Phe Ile Leu
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Input Set : A:\CANDIDA.txt

Output Set: N:\CRF3\03022001\I763011.raw

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368 Ser Glu Leu Gly Thr Lys Thr Pro Asn Phe Glu Asp Pro Phe Glu Gln
370                               370                               375                               380
374 Ile Asn Ile Glu Asp Asn Lys Val Ile Ser Val Asn Thr Pro Lys Thr
376 385                               390                               395                               400
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382                               405                               410                               415
386 Thr Phe Glu Ser Thr Ile Phe Glu Ile Pro Lys Phe Phe Tyr Gly Ser
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394                               435                               440                               445
398 Phe Asp Val Phe Asp Trp Ile Phe Glu Ser Gly Thr Thr Asn Glu Lys
400                               450                               455                               460
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488 acgttatcta caccaatgaa caaaaatcct gttcgtggac atagtaccaa gaaaaagctt 1800
490 agtatgttgg acatgcattg gtctttctaaa attcaaagtc ttttacctcc acaaccgcca 1860
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494 ttgacttatac aaagagaatt tttcaattat aaaaaagtga ttgttcaata ccaaatggaa 1980
496 cggataaaca aagaccttga acattttgac gatataaatg atggttcaca cactgagaat 2040
498 ttggataactt tcaagcattg tttagaacaa gattatttgg ttatgagtga gtttaatgaa 2100
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526 20 25 30
530 Tyr His Pro Asp Lys Thr Pro Arg Arg Glu Asp His Glu Lys Phe Lys
532 35 40 45
536 Glu Ile Asn Ile Ala Tyr Glu Thr Ile Arg Asp Tyr Tyr Gln Glu Asn
538 50 55 60
542 Gly Gln Lys Asn Ser Gln Pro Ile Pro Asn Thr Asn Thr Glu His Asn
544 65 70 75 80
548 Ser His Gln Lys Pro His Tyr Asn Thr Gly Pro Tyr Ser Thr Tyr Arg
550 85 90 95
554 Phe Thr Thr Ser Ser Thr Thr Thr Asp Asn Thr Asn His Thr Gly His
556 100 105 110
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562 115 120 125
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568 130 135 140
572 Lys Lys Glu Leu Phe Gln Arg Gln Gln Ala Glu Glu Ala Gln Arg Lys
574 145 150 155 160
578 Lys Glu Phe Glu Gln Lys Ala Glu Phe Ile Lys Ala Ser Leu Leu Glu
580 165 170 175
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586 180 185 190
590 Gln Arg Gln Lys Glu His Glu Ala Lys Arg Asp Ile Arg Ile Gln Gln
592 195 200 205
596 Leu Ser Glu Gln Asp Ser Arg Ser Asn Gln Thr Lys Glu Glu Glu Glu
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<211> 302

<212> PRT

<213> Candida albicans

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10

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Gly Asn Pro Val Ala Val Ile Tyr Asp Ser Asp Asn Leu Thr Thr Gln

20

25

30

Glu Met Gln Lys Ile Ala Arg Trp Thr Asn Leu Ser Glu Thr Thr Phe

35

40

45

Ile Leu Thr Pro Lys Ser Ser Ile Ala Xaa Tyr Ser Ile Arg Ile Phe

50

55

60

Thr Ser Gly Gly Asn Glu Leu Pro Phe Ala Gly His Pro Thr Leu Gly

65

70

75

80

Thr Ala Phe Ala Leu Leu Glu Asp Gly Lys Ile Lys Pro Asn Asp Asn

85

90

95

Gly Gln Ile Ile Gln Glu Cys Gly Ala Gly Leu Val Lys Ile Ser Val

100

105

110

Glu Lys Thr Pro Asn Asn Asn Ser Asn Glu Leu Pro Phe Leu Leu Ser

115

120

125

→ see
item 10
on Enon
Summary
Sheet

09/763,011

7

Phe Glu Leu Pro Tyr Phe Lys Phe His Glu Ile Asp Asp Lys Val Ile
 130 135 140

Glu Glu Leu Gln His Ser Trp Asn Gly Thr Asn Ile Ile Gly Lys Pro
 145 150 155 160

Val Leu Ile Asp Ala Gly Pro Lys Trp Ala Val Phe Gln Leu Gly Ser
 165 170 175

Gly Lys Glu Val Leu Asp Leu Asn (Xaa) Asp Leu Ala Gln Ile Glu Arg
 180 185 190

Leu Ser Leu Glu Asn Gly Trp Thr Gly Ile Gly Val Phe Gly Lys His
 195 200 205

Asn Glu Asn Gly Asp Ser Val Glu Leu Arg Asn Ile Ala Pro Ala Val
 210 215 220

Gly Val Ala Glu Asp Pro Ala Cys Gly Ser Gly Ser Gly Ala Ile Gly
 225 230 235 240

Ala Tyr Leu Ala Asn His Val Phe Asn Glu Lys Glu Lys Phe Thr Ile
 245 250 255

Asp Ile Ser Gln Gly Lys Pro Ile Glu Arg Asp Ala Lys Ile Gln Val
 260 265 270

only
 covered
 portions
 of
 seq. 13
 shown

see item 10

FYI

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

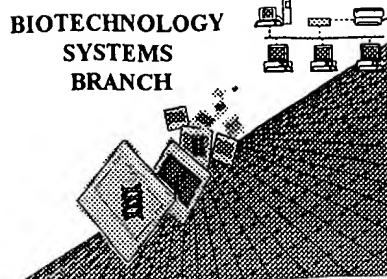
VERIFICATION SUMMARY
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L:19 M:270 C: Current Application Number differs, Replaced Application Number
L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1060 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:13
L:1060 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:13
L:1060 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
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L:1060 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:13
L:1108 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:13
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M:340 Repeated in SeqNo=13
L:5730 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:57
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M:340 Repeated in SeqNo=114

RAW SEQUENCE LISTING **ERROR REPORT**



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/763,011

Source: Pt 09

Date Processed by STIC: 3/2/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>